WQ 01/49826



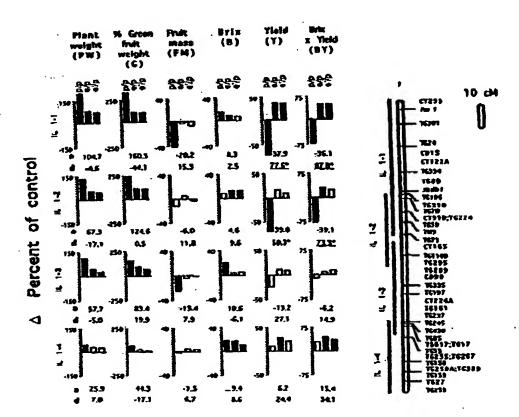


Fig. 1a (Prior Art)

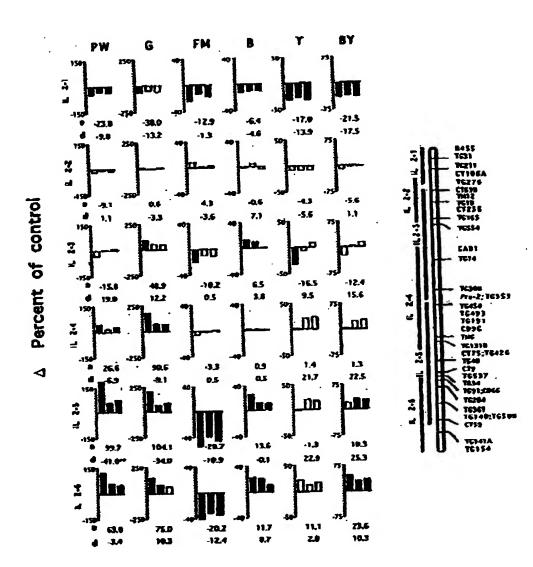
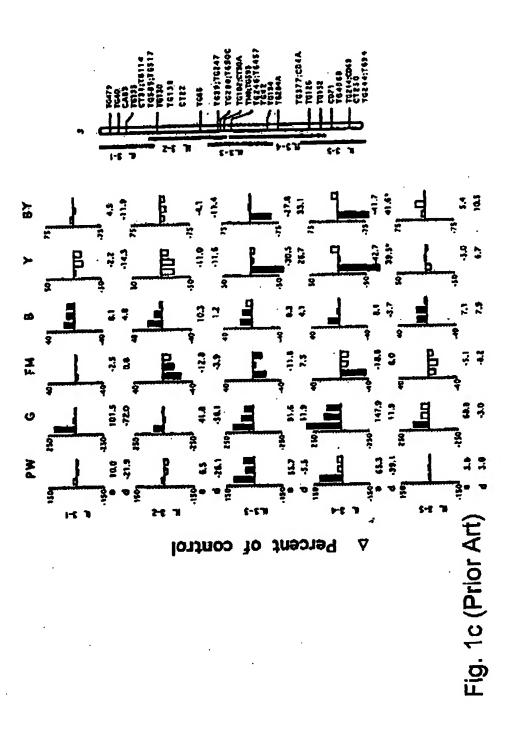


Fig. 1b (Prior Art)

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PT/JL01/000 0 / 0 7 0 9 2 3



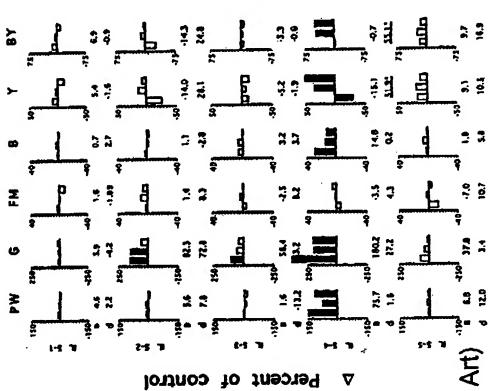
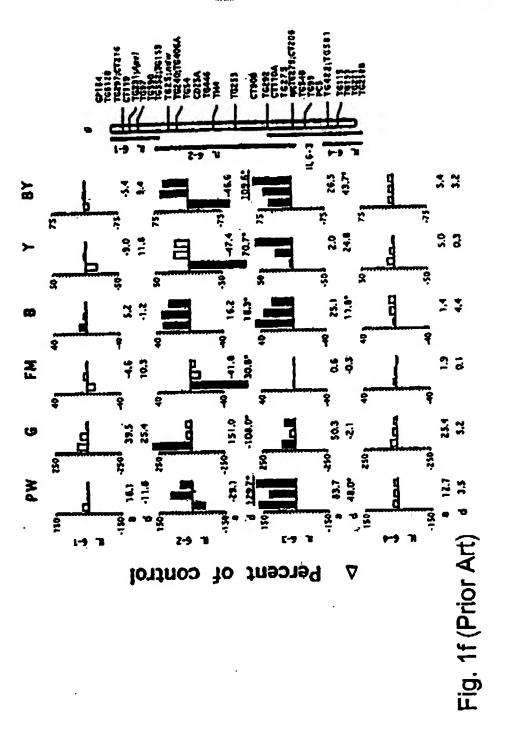


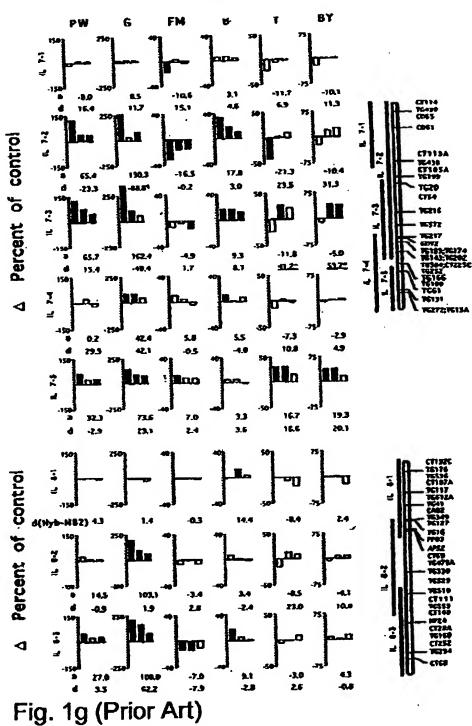
Fig. 1e (Prior Art)

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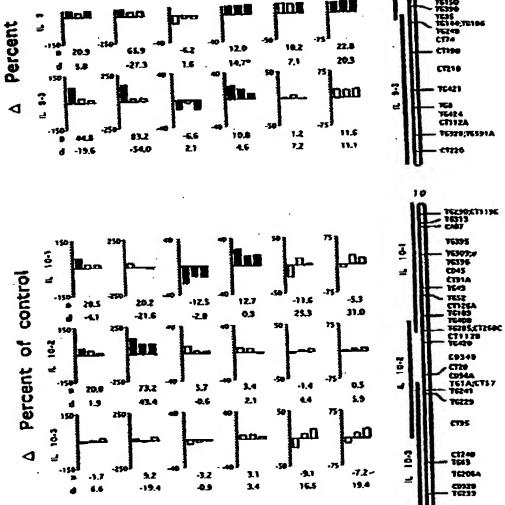


Fig. 1h (Prior Art)

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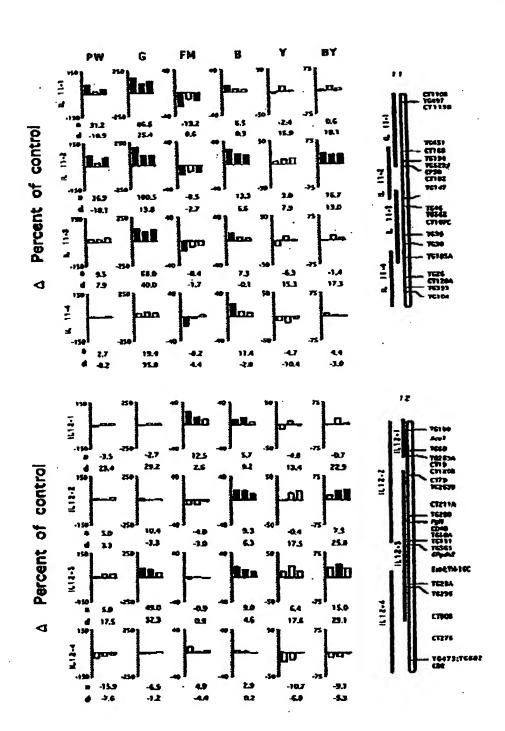


Fig. 1i (Prior Art)

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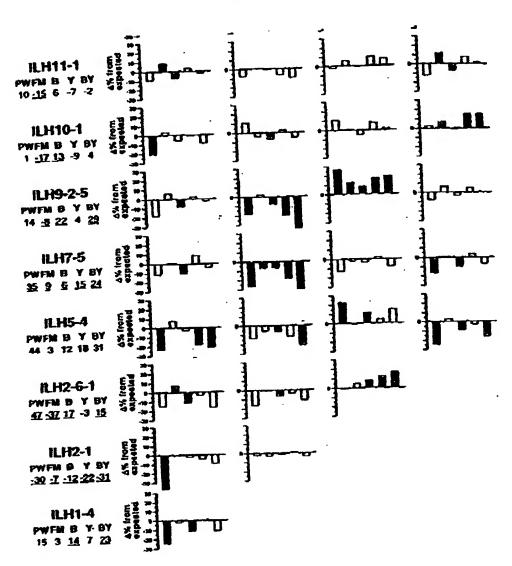


Fig. 2 (Prior Art)

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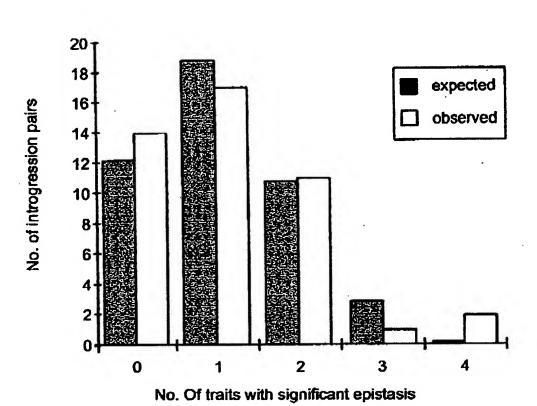
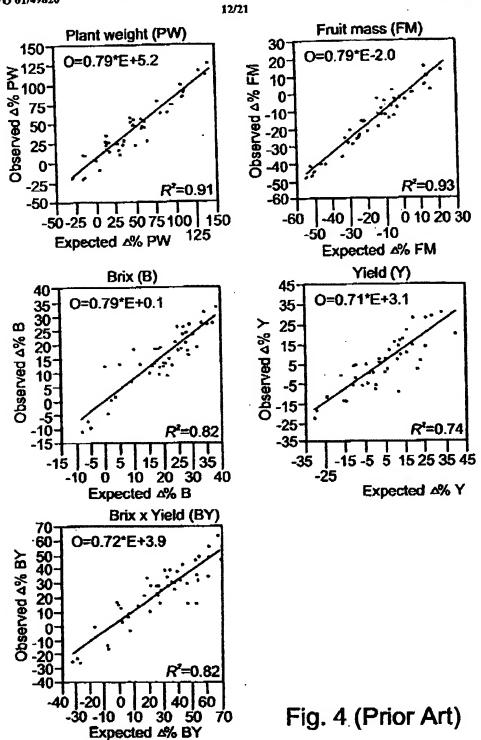


Fig. 3 (Prior Art)

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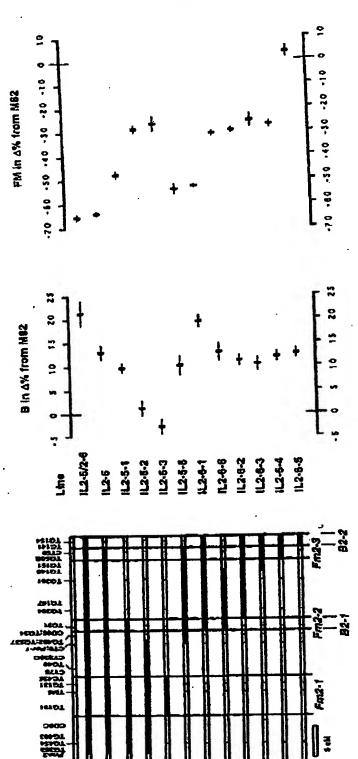
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Fig. 5 (Prior Art)

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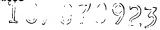




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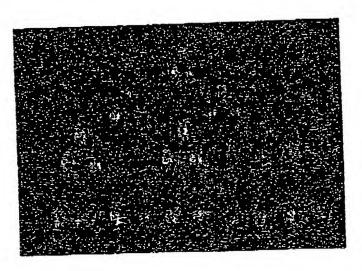
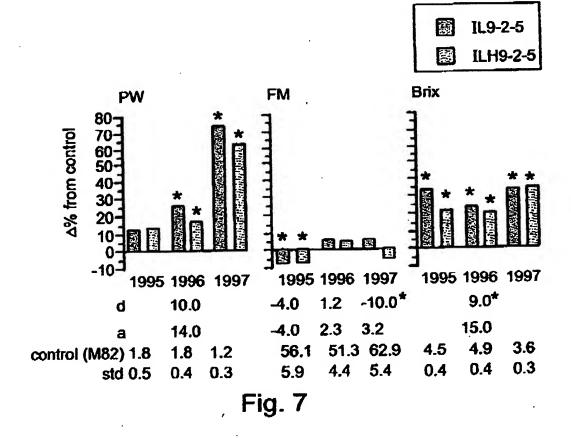


Fig. 6 (Prior Art)



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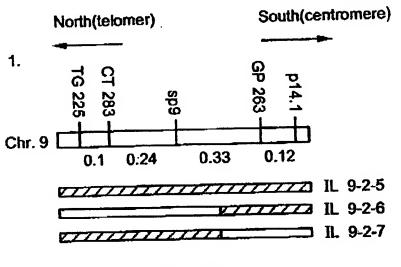
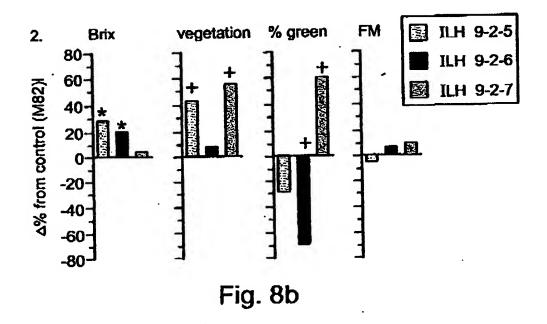
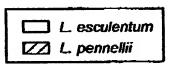
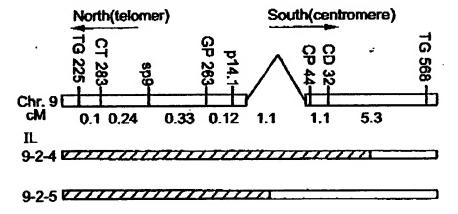


Fig. 8a



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-IL 9-2-5 X 202

-M82 X 202

Fig. 10

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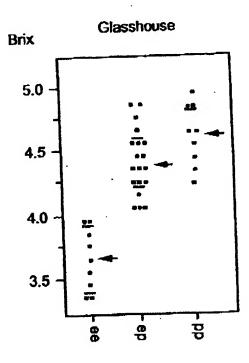
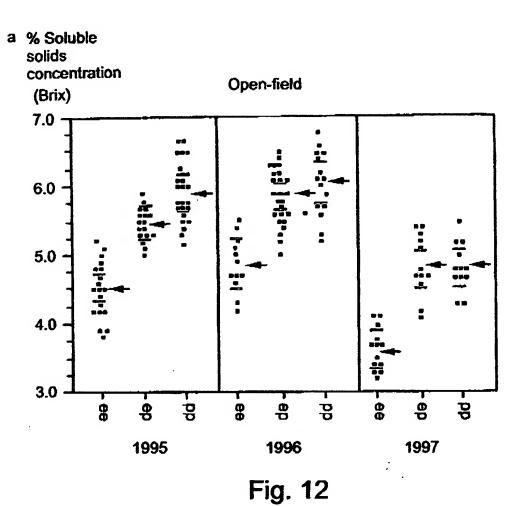


Fig. 11

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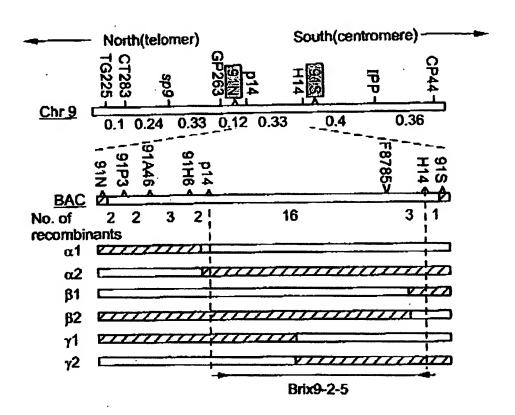


Fig. 13

TL01/00008 0/070923

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TT CG AT CAAG TA CC TTGC SNP 2963 2935 Phenotypic effect (%) Position 2 +10+20 +30 Rec. 8 10 43 13 6 29 29 Brix9-2-5 100 bp 29

20/21

100 bp

10 17 4 6

22 1 1 13

23 1 13

25

Exons I

III

IV

V

VI

L. pennellii Brix9-2-5 allele

Fig. 14a

2301 GACINIGGIA ATTICIANGC ATCAMBACA TICTATGATC CIAGCAGNAA
2351 TCGAAGGGIT ATTIGGGGIT GGICAANTGA ATCAGATGIA TERCCIGACG
2401 ATGATATINA GAAAGGATGG CCTGGANITC AAGGIATICC GCGACAAGTA
2451 TGGCTAGACC TIMGIGGIAN ACARTRAGIT CAATGGCCTA TIGAAGAATT
2501 AGAAACCCTA AGGAAGCANA AGGICCAATT GAACAACAAG AAGTIGAGCA
2551 ACGGAGAAAT GITTGAAGIT AAAGGANTCI CAGCATCACA GGITTGAACT
2601 TTIACTIAIT AACCCAINGI CITITAAATA TCATTAATCI AGTICTIAIG
2651 TATACAGTGI ATAAGCAATA TATACTAATT
2701 ATCCATATAT ATAAGTAGA FIGAATTATAC ATTIGITATA TATACTAATT
2751 ATAACAATG CACTGICCCA TCTTGCATTT CTTTGATTGI AGGCTGATGI
2801 TGAAGTGITA TTCTCATTTT CAAGTTAAA CAAGGCCGAA CAATTGATC

Fig. 14b

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